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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/972,970

DATE: 10/19/2001
 TIME: 12:12:03

Input Set : A:\PT056.seqlist.txt
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2 <110> APPLICANT: Shi et al.
 4 <120> TITLE OF INVENTION: TM4SF Receptor Polynucleotides, Polypeptides, and Antibodies
 6 <130> FILE REFERENCE: PT056P1
 C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/972,970
 9 <141> CURRENT FILING DATE: 2001-10-10
 11 <150> PRIOR APPLICATION NUMBER: PCT/US01/11130
 12 <151> PRIOR FILING DATE: 2001-04-05
 14 <150> PRIOR APPLICATION NUMBER: 60/195,336
 15 <151> PRIOR FILING DATE: 2000-04-10
 17 <160> NUMBER OF SEQ ID NOS: 8
 19 <170> SOFTWARE: PatentIn Ver. 2.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 733
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Homo sapiens
 27 <400> SEQUENCE: 1

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30	tctcccgac	tcctgagggtc	acatgcgtgg	tggtgacgt	aagccacgaa	gaccctgagg	180
31	tcaagttcaa	ctggtaacgtg	gacggcgtgg	aggtgcataa	tgccaagaca	aagccgcggg	240
32	aggagcagta	caacagcacg	taccgtgtgg	tcagcgtcct	caccgtcctg	caccaggact	300
33	ggctgaatgg	caaggagtac	aagtgcagg	tctccaaacaa	agccctccca	accccccattcg	360
34	agaaaaccat	ctccaaagcc	aaagggcagc	cccgagaacc	acaggtgtac	accctgcccc	420
35	catcccgaaa	tgagctgacc	aagaaccagg	tcagcgtgac	ctgcctggtc	aaaggcttct	480
36	atccaagcga	catcgccgtg	gagtggaga	gcaatggca	gccggagaaac	aactacaaga	540
37	ccacgcctcc	cgtgctggac	tccgacggct	ccttcttcct	ctacagcaag	ctcaccgtgg	600
38	acaagagcag	gtggcagcag	gggaacgtct	tctcatgctc	cgtgatgcat	gaggctctgc	660
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51	gccgggcaag	caccagca	tccaggaacc	cgaggtcggc	tgctgcggga	aataacttcct	180
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59	taatgactgg	aacctaata	tctatttcaa	ctgcaactgac	ttgaacccga	gccgagagcg	660
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64	atcaaggcag tgaagccaa ctggatcaa catgatgatg gctacaaact actcaaataa	960
65	acaaaacctt gaaaaccact gccttacgcc caccatctca gaggttccat gggccgcagg	1020
66	gcctcagccg tgccgtctgc ctggggcccc agcccagacc caccctgcca acattttc	1080
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77	gtctgcctt cctaggacaa ccccacccag tacagccctg tgcctgggt gtccaccctg	1740
78	cttactagtt ctgtgggtt catggattt acaagtttca aaggagcag agtgcgtcag	1800
79	attggggaaag cctggcagct gttctcagat ctgcacaaag cggtgtgtgt ggagtatttg	1860
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82	tcccactatt cctggagatg ggacatagag agacattgtt ttgtgctgag aaataacttc	2040
83	atgattgagt ctgagtcgtc aaggcaact ggccttgagt gacatcaagg ggtgtgggg	2100
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89	gaactgctgc aggtggccgg ttgtgggccc agcaccatc gtggctccca ggtgtggct	2460
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102	ttttccttct tctacatcct tgattatttc cttaggtcca gttttataa gtcaattac	180
103	atggtcaaag ggatgttggaa ttgggtgtt gctttgata ttactaaatt gatcttcaca	240
104	aaggattcac cagtccaaagc tcttacttgtt gatgtcaat gtgttgcgt tactgtatga	300
105	tctctactcc taggattgtt tacaacaa gtcagaaacc aacaaataca agcctgtcat	360
106	attttcgat ttgttggatg gaaattatct tttcttctac aattttctga tttctactt	420
107	tttacattaa ctttcgttga taaaccttct ttatgttct ttattatagg aaattatagg	480
108	agatttattt agtttcttca tggctgtgtt tagcggctct aggttagatcg gtataatcat	540
109	aaaaatgaaa tagtactta aaattggcca ttgttgcac gtagcaggca ttatgttgg	600
110	tgctttacaa aatatacgatc atttaatccc cgccacaaatt ctggaggtag gaaatattat	660
111	tctcatatca ctgagaaatgtt attgttgc accatttattt attcacctt gtgttgcgt	720
112	agtttggcca agtacactggc acattcacac aaattgggtt ttgttgcacatgaa cattagagaa	780
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116	ccccaaatggc	taaccaccaa	tccctaaaat	tgaatttttg	ttctgttggg	tgtcctctga	1020
117	gttcgtgtct	atggttacac	aatacccaag	attataaaaa	ctgaaaaatg	ttgtcagtgt	1080
118	ccaaacaaacc	aattcaaata	aatgatatcc	aaatatcata	tagacattaa	cctctttatg	1140
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120	aaaatttgaac	agcatttca	gatcgaattt	tctgatffff	gcactttgct	gcttgggtcc	1260
121	aaataaaaaat	agaacagaat	gaagtggatc	aaaagccaaa	tcaaaaat	ttttccctt	1320
122	gtgatattt	aatttagtac	tacaaagcat	tgaaatgaga	gatgctagta	aatggatat	1380
123	atgttattat	tgcttggct	ttgtcagttt	tatgacctt	cttggctt	cagatttt	1440
124	tcttacagt	tattcagaat	tatgtgaatt	aggattctct	ttaatgtaga	atgcaattt	1500
125	aattattggc	ttaatagctt	aaaatgaaca	gtcttaaaca	gtcttgcaaa	ttctttgtct	1560
126	tggaaagctgg	gaactgttca	atctctggaa	cagtggtcat	aggatagtct	cattaatcat	1620
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143	35	40	45				
145	Leu Ser Asn Ile Ser Ala Leu Thr Asp Leu Gly Gly Leu Asp Pro Val						
146	50	55	60				
148	Trp Leu Phe Val Val Val Gly Gly Val Met Ser Val Leu Gly Phe Ala						
149	65	70	75	80			
151	Gly Cys Ile Gly Ala Leu Arg Glu Asn Thr Phe Leu Leu Lys Phe Phe						
152	85	90	95				
154	Ser Val Phe Leu Gly Leu Ile Phe Phe Leu Glu Leu Ala Thr Gly Ile						
155	100	105	110				
157	Leu Ala Phe Val Phe Lys Asp Trp Ile Arg Asp Gln Leu Asn Phe Phe						
158	115	120	125				
160	Ile Asn Asn Asn Val Lys Ala Tyr Arg Asp Asp Ile Asp Leu Gln Asn						
161	130	135	140				
163	Leu Ile Asp Phe Ala Gln Glu Tyr Trp Ser Cys Cys Gly Ala Arg Gly						
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166	Pro Asn Asp Trp Asn Leu Asn Ile Tyr Phe Asn Cys Thr Asp Leu Asn						
167	165	170	175				
169	Pro Ser Arg Glu Arg Cys Gly Val Pro Phe Ser Cys Cys Val Arg Asp						
170	180	185	190				
172	Pro Ala Met Ser Ser Thr Pro Ser Val Ala Met Met Ser Gly Ser Asn						
173	195	200	205				
175	Trp Ser Trp Ser Ser Arg Ala Pro Tyr Thr Pro Lys Ala Val Trp Ala						
176	210	215	220				
178	Ser Leu Arg Ser Gly Cys Arg Thr Thr						
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183 <211> LENGTH: 44
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191 Val Ser Phe Met Thr Phe Leu Trp Leu Ser Asp Leu Phe Leu Thr Val
192 20 25 30
194 Tyr Ser Glu Leu Cys Glu Leu Gly Phe Ser Leu Met
195 35 40
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199 <211> LENGTH: 18
200 <212> TYPE: PRT
201 <213> ORGANISM: Homo sapiens
203 <400> SEQUENCE: 6
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205 1 5 10 15
207 Ile Gly
211 <210> SEQ ID NO: 7
212 <211> LENGTH: 17
213 <212> TYPE: PRT
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225 <211> LENGTH: 20
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229 <400> SEQUENCE: 8
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VERIFICATION SUMMARY

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Input Set : A:\PT056.seqlist.txt

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L:8 M:270 C: Current Application Number differs, Replaced Current Application Number